

RECEIVED

CRF Errors Corrected by the STIC Systems Branch.

CRF Processing Date: 5/28/2003Edited by: Ar

Verified by: _____

(STIC staff)

TECH CENTER 1600/2900

Serial Number: 09/919,408A☐

Changed a file from non-ASCII to ASCII

☐

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐

Edited a format error in the Current Application Data section, specifically: _____

☐Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____☐

Added the mandatory heading and subheadings for "Current Application Data".

☐

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐

Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐

Inserted colons after headings/subheadings. Headings edited included: _____

☐

Deleted extra, invalid, headings used by an applicant, specifically: _____

☐Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____☐

Inserted mandatory headings, specifically: _____

☐

Corrected an obvious error in the response, specifically: _____

☐

Edited identifiers where upper case is used but lower case is required, or vice versa.

☐

Corrected an error in the Number of Sequences field, specifically: _____

☐

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____☒Other: Sequence 11 - aligned amino acid nos.

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

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JUN 04 2003

TECH CENTER



RAW SEQUENCE LISTING

DATE: 05/28/2003

PATENT APPLICATION: US/09/919,408A

TIME: 11:31:13

Input Set : N:\Cr4\05272003\I919408A.raw

Output Set: N:\CRF4\05282003\I919408A.raw

SEQUENCE LISTING

1 (1) GENERAL INFORMATION:

2 (i) APPLICANT: Lemischka, Ihor R.

 3 (ii) TITLE OF INVENTION: METHOD FOR ISOLATING CELLS EXPRESSING
 4 FLK-2 RECEPTORS AND ISOLATED POPULATIONS
 5 OF CELLS THAT EXPRESS FLK-2 RECEPTORS

6 (iii) NUMBER OF SEQUENCES: 11

7 (iv) CORRESPONDENCE ADDRESS:

8 (A) ADDRESSEE: Kenyon & Kenyon

9 (B) STREET: One Broadway

10 (C) CITY: New York

11 (D) STATE: New York

12 (E) COUNTRY: U.S.A.

13 (F) ZIP: 10004

14 (v) COMPUTER READABLE FORM:

15 (A) MEDIUM TYPE: Floppy disk

16 (B) COMPUTER: IBM PC compatible

17 (C) OPERATING SYSTEM: PC-DOS/MS-DOS

18 (D) SOFTWARE: PatentIn Release #1.0, Version #1.25

19 (vi) CURRENT APPLICATION DATA:

C--> 20 (A) APPLICATION NUMBER: US/09/919,408A

C--> 21 (B) FILING DATE: 31-Jul-2001

22 (vii) PRIOR APPLICATION DATA:

W--> 23 (A) APPLICATION NUMBER: US 09/208,786

24 (B) FILING DATE: 10-DEC-1998

W--> 25 (A) APPLICATION NUMBER: US 09/021,324

26 (B) FILING DATE: 10-FEB-1998

W--> 27 (A) APPLICATION NUMBER: US 08/601,891

28 (B) FILING DATE: 15-FEB-1996

W--> 29 (A) APPLICATION NUMBER: US 08/252,498

30 (B) FILING DATE: 31-OCT-1994

W--> 31 (A) APPLICATION NUMBER: US 08/055,269

32 (B) FILING DATE: 30-APR-1993

W--> 33 (A) APPLICATION NUMBER: US 07/977,451

34 (B) FILING DATE: 19-NOV-1992

W--> 35 (A) APPLICATION NUMBER: US 07/975,049

36 (B) FILING DATE: 12-NOV-1992

W--> 37 (A) APPLICATION NUMBER: 07/906,397

38 (B) FILING DATE: 26-JUN-1992

W--> 39 (A) APPLICATION NUMBER: 07/813,593

40 (B) FILING DATE: 24-DEC-1991

W--> 41 (A) APPLICATION NUMBER: 07/793,065

42 (B) FILING DATE: 15-NOV-1991

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Input Set : N:\Cr4\05272003\I919408A.raw

Output Set: N:\CRF4\05282003\I919408A.raw

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W--> 43      (A) APPLICATION NUMBER: 07/728,913
44          (B) FILING DATE: 28-JUN-1991
W--> 45      (A) APPLICATION NUMBER: 07/679,666
46          (B) FILING DATE: 02-APR-1991
47      (viii) ATTORNEY/AGENT INFORMATION:
48          (A) NAME: Wieckowski, Elizabeth M.
49          (B) REGISTRATION NUMBER: 42,226
50          (C) REFERENCE/DOCKET NUMBER: 11245/46115
51      (ix) TELECOMMUNICATION INFORMATION:
52          (A) TELEPHONE: 212-425-7200
53          (B) TELEFAX: 212-425-5288
54      (2) INFORMATION FOR SEQ ID NO: 1:
55          (i) SEQUENCE CHARACTERISTICS:
56              (A) LENGTH: 3453 base pairs
57              (B) TYPE: nucleic acid
58              (C) STRANDEDNESS: double
59              (D) TOPOLOGY: linear
60          (ii) MOLECULE TYPE: cDNA
61          (iii) HYPOTHETICAL: NO
62          (iv) ANTI-SENSE: NO
63          (v) FRAGMENT TYPE: N-terminal
64          (ix) FEATURE:
65              (A) NAME/KEY: mat_peptide
66              (B) LOCATION: 112..3006
67          (ix) FEATURE:
68              (A) NAME/KEY: sig_peptide
69              (B) LOCATION: 31..111
70          (ix) FEATURE:
71              (A) NAME/KEY: CDS
72              (B) LOCATION: 31..3009
73          (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
74      GCGGCCTGGC TACCGCGCGC TCCGGAGGCC ATG CGG GCG TTG GCG CAG CGC AGC      54
75                                     Met Arg Ala Leu Ala Gln Arg Ser
76                                     -27   -25   -20
77      GAC CGG CGG CTG CTG CTG CTT GTT GTT TTG TCA GTA ATG ATT CTT GAG      102
78      Asp Arg Arg Leu Leu Leu Leu Val Val Leu Ser Val Met Ile Leu Glu
79                                     -15   -10   -5
80      ACC GTT ACA AAC CAA GAC CTG CCT GTG ATC AAG TGT GTT TTA ATC AGT      150
81      Thr Val Thr Asn Gln Asp Leu Pro Val Ile Lys Cys Val Leu Ile Ser
82                                     1       5       10
83      CAT GAG AAC AAT GGC TCA TCA GCG GGA AAG CCA TCA TCG TAC CGA ATG      198
84      His Glu Asn Asn Gly Ser Ser Ala Gly Lys Pro Ser Ser Tyr Arg Met
85      15       20       25
86      GTG CGA GGA TCC CCA GAA GAC CTC CAG TGT ACC CCG AGG CGC CAG AGT      246
87      Val Arg Gly Ser Pro Glu Asp Leu Gln Cys Thr Pro Arg Arg Gln Ser
88      30       35       40       45
89      GAA GGG ACG GTA TAT GAA GCG GCC ACC GTG GAG GTG GCC GAG TCT GGG      294
90      Glu Gly Thr Val Tyr Glu Ala Ala Thr Val Glu Val Ala Glu Ser Gly
91                                     50       55       60

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92	TCC ATC ACC CTG CAA GTG CAG CTC GCC ACC CCA GGG GAC CTT TCC TGC	342
93	Ser Ile Thr Leu Gln Val Gln Leu Ala Thr Pro Gly Asp Leu Ser Cys	
94	65 70 75	
95	CTC TGG GTC TTT AAG CAC AGC TCC CTG GGC TGC CAG CCG CAC TTT GAT	390
96	Leu Trp Val Phe Lys His Ser Ser Leu Gly Cys Gln Pro His Phe Asp	
97	80 85 90	
98	TTA CAA AAC AGA GGA ATC GTT TCC ATG GCC ATC TTG AAC GTG ACA GAG	438
99	Leu Gln Asn Arg Gly Ile Val Ser Met Ala Ile Leu Asn Val Thr Glu	
100	95 100 105	
101	ACC CAG GCA GGA GAA TAC CTA CTC CAT ATT CAG AGC GAA CGC GCC AAC	486
102	Thr Gln Ala Gly Glu Tyr Leu Leu His Ile Gln Ser Glu Arg Ala Asn	
103	110 115 120 125	
104	TAC ACA GTA CTG TTC ACA GTG AAT GTA AGA GAT ACA CAG CTG TAT GTG	534
105	Tyr Thr Val Leu Phe Thr Val Asn Val Arg Asp Thr Gln Leu Tyr Val	
106	130 135 140	
107	CTA AGG AGA CCT TAC TTT AGG AAG ATG GAA AAC CAG GAT GCA CTG CTC	582
108	Leu Arg Arg Pro Tyr Phe Arg Lys Met Glu Asn Gln Asp Ala Leu Leu	
109	145 150 155	
110	TGC ATC TCC GAG GGT GTT CCG GAG CCC ACT GTG GAG TGG GTG CTC TGC	630
111	Cys Ile Ser Glu Gly Val Pro Glu Pro Thr Val Glu Trp Val Leu Cys	
112	160 165 170	
113	AGC TCC CAC AGG GAA AGC TGT AAA GAA GAA GGC CCT GCT GTT GTC AGA	678
114	Ser Ser His Arg Glu Ser Cys Lys Glu Glu Gly Pro Ala Val Val Arg	
115	175 180 185	
116	AAG GAG GAA AAG GTA CTT CAT GAG TTG TTC GGA ACA GAC ATC AGA TGC	726
117	Lys Glu Glu Lys Val Leu His Glu Leu Phe Gly Thr Asp Ile Arg Cys	
118	190 195 200 205	
119	TGT GCT AGA AAT GCA CTG GGC CGC GAA TGC ACC AAG CTG TTC ACC ATA	774
120	Cys Ala Arg Asn Ala Leu Gly Arg Glu Cys Thr Lys Leu Phe Thr Ile	
121	210 215 220	
122	GAT CTA AAC CAG GCT CCT CAG AGC ACA CTG CCC CAG TTA TTC CTG AAA	822
123	Asp Leu Asn Gln Ala Pro Gln Ser Thr Leu Pro Gln Leu Phe Leu Lys	
124	225 230 235	
125	GTG GGG GAA CCC TTG TGG ATC AGG TGT AAG GCC ATC CAT GTG AAC CAT	870
126	Val Gly Glu Pro Leu Trp Ile Arg Cys Lys Ala Ile His Val Asn His	
127	240 245 250	
128	GGA TTC GGG CTC ACC TGG GAG CTG GAA GAC AAA GCC CTG GAG GAG GGC	918
129	Gly Phe Gly Leu Thr Trp Glu Leu Glu Asp Lys Ala Leu Glu Glu Gly	
130	255 260 265	
131	AGC TAC TTT GAG ATG AGT ACC TAC TCC ACA AAC AGG ACC ATG ATT CGG	966
132	Ser Tyr Phe Glu Met Ser Thr Tyr Ser Thr Asn Arg Thr Met Ile Arg	
133	270 275 280 285	
134	ATT CTC TTG GCC TTT GTG TCT TCC GTG GGA AGG AAC GAC ACC GGA TAT	1014
135	Ile Leu Leu Ala Phe Val Ser Ser Val Gly Arg Asn Asp Thr Gly Tyr	
136	290 295 300	
137	TAC ACC TGC TCT TCC TCA AAG CAC CCC AGC CAG TCA GCG TTG GTG ACC	1062
138	Tyr Thr Cys Ser Ser Ser Lys His Pro Ser Gln Ser Ala Leu Val Thr	
139	305 310 315	
140	ATC CTA GAA AAA GGG TTT ATA AAC GCT ACC AGC TCG CAA GAA GAG TAT	1110

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141	Ile	Leu	Glu	Lys	Gly	Phe	Ile	Asn	Ala	Thr	Ser	Ser	Gln	Glu	Glu	Tyr	
142			320					325					330				
143	GAA	ATT	GAC	CCG	TAC	GAA	AAG	TTC	TGC	TTC	TCA	GTC	AGG	TTT	AAA	GCG	1158
144	Glu	Ile	Asp	Pro	Tyr	Glu	Lys	Phe	Cys	Phe	Ser	Val	Arg	Phe	Lys	Ala	
145			335					340					345				
146	TAC	CCA	CGA	ATC	CGA	TGC	ACG	TGG	ATC	TTC	TCT	CAA	GCC	TCA	TTT	CCT	1206
147	Tyr	Pro	Arg	Ile	Arg	Cys	Thr	Trp	Ile	Phe	Ser	Gln	Ala	Ser	Phe	Pro	
148			350					355				360				365	
149	TGT	GAA	CAG	AGA	GGC	CTG	GAG	GAT	GGG	TAC	AGC	ATA	TCT	AAA	TTT	TGC	1254
150	Cys	Glu	Gln	Arg	Gly	Leu	Glu	Asp	Gly	Tyr	Ser	Ile	Ser	Lys	Phe	Cys	
151					370					375					380		
152	GAT	CAT	AAG	AAC	AAG	CCA	GGA	GAG	TAC	ATA	TTC	TAT	GCA	GAA	AAT	GAT	1302
153	Asp	His	Lys	Asn	Lys	Pro	Gly	Glu	Tyr	Ile	Phe	Tyr	Ala	Glu	Asn	Asp	
154				385					390					395			
155	GAC	GCC	CAG	TTC	ACC	AAA	ATG	TTC	ACG	CTG	AAT	ATA	AGA	AAG	AAA	CCT	1350
156	Asp	Ala	Gln	Phe	Thr	Lys	Met	Phe	Thr	Leu	Asn	Ile	Arg	Lys	Lys	Pro	
157			400					405					410				
158	CAA	GTG	CTA	GCA	AAT	GCC	TCA	GCC	AGC	CAG	GCG	TCC	TGT	TCC	TCT	GAT	1398
159	Gln	Val	Leu	Ala	Asn	Ala	Ser	Ala	Ser	Gln	Ala	Ser	Cys	Ser	Ser	Asp	
160			415					420					425				
161	GGC	TAC	CCG	CTA	CCC	TCT	TGG	ACC	TGG	AAG	AAG	TGT	TCG	GAC	AAA	TCT	1446
162	Gly	Tyr	Pro	Leu	Pro	Ser	Trp	Thr	Trp	Lys	Lys	Cys	Ser	Asp	Lys	Ser	
163			430					435				440				445	
164	CCC	AAT	TGC	ACG	GAG	GAA	ATC	CCA	GAA	GGA	GTT	TGG	AAT	AAA	AAG	GCT	1494
165	Pro	Asn	Cys	Thr	Glu	Ile	Pro	Glu	Gly	Val	Trp	Asn	Lys	Lys	Ala		
166					450					455					460		
167	AAC	AGA	AAA	GTG	TTT	GGC	CAG	TGG	GTG	TCG	AGC	AGT	ACT	CTA	AAT	ATG	1542
168	Asn	Arg	Lys	Val	Phe	Gly	Gln	Trp	Val	Ser	Ser	Ser	Thr	Leu	Asn	Met	
169				465					470					475			
170	AGT	GAG	GCC	GGG	AAA	GGG	CTT	CTG	GTC	AAA	TGC	TGT	GCG	TAC	AAT	TCT	1590
171	Ser	Glu	Ala	Gly	Lys	Gly	Leu	Leu	Val	Lys	Cys	Cys	Ala	Tyr	Asn	Ser	
172			480					485					490				
173	ATG	GGC	ACG	TCT	TGC	GAA	ACC	ATC	TTT	TTA	AAC	TCA	CCA	GGC	CCC	TTC	1638
174	Met	Gly	Thr	Ser	Cys	Glu	Thr	Ile	Phe	Leu	Asn	Ser	Pro	Gly	Pro	Phe	
175			495					500				505					
176	CCT	TTC	ATC	CAA	GAC	AAC	ATC	TCC	TTC	TAT	GCG	ACC	ATT	GGG	CTC	TGT	1686
177	Pro	Phe	Ile	Gln	Asp	Asn	Ile	Ser	Phe	Tyr	Ala	Thr	Ile	Gly	Leu	Cys	
178			510					515				520				525	
179	CTC	CCC	TTC	ATT	GTT	GTT	CTC	ATT	GTG	TTG	ATC	TGC	CAC	AAA	TAC	AAA	1734
180	Leu	Pro	Phe	Ile	Val	Leu	Ile	Val	Leu	Ile	Cys	His	Lys	Tyr	Lys		
181					530					535					540		
182	AAG	CAA	TTT	AGG	TAC	GAG	AGT	CAG	CTG	CAG	ATG	ATC	CAG	GTG	ACT	GGC	1782
183	Lys	Gln	Phe	Arg	Tyr	Glu	Ser	Gln	Leu	Gln	Met	Ile	Gln	Val	Thr	Gly	
184				545					550					555			
185	CCC	CTG	GAT	AAC	GAG	TAC	TTC	TAC	GTT	GAC	TTC	AGG	GAC	TAT	GAA	TAT	1830
186	Pro	Leu	Asp	Asn	Glu	Tyr	Phe	Tyr	Val	Asp	Phe	Arg	Asp	Tyr	Glu	Tyr	
187			560					565					570				
188	GAC	CTT	AAG	TGG	GAG	TTC	CCG	AGA	GAG	AAC	TTA	GAG	TTT	GGG	AAG	GTC	1878
189	Asp	Leu	Lys	Trp	Glu	Phe	Pro	Arg	Glu	Asn	Leu	Glu	Phe	Gly	Lys	Val	

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190	575	580	585	
191	CTG GGG TCT GGC GCT TTC GGG AGG GTG ATG AAC GCC ACG GCC TAT GGC	1926		
192	Leu Gly Ser Gly Ala Phe Gly Arg Val Met Asn Ala Thr Ala Tyr Gly			
193	590 595 600 605			
194	ATT AGT AAA ACG GGA GTC TCA ATT CAG GTG GCG GTG AAG ATG CTA AAA	1974		
195	Ile Ser Lys Thr Gly Val Ser Ile Gln Val Ala Val Lys Met Leu Lys			
196	610 615 620			
197	GAG AAA GCT GAC AGC TGT GAA AAA GAA GCT CTC ATG TCG GAG CTC AAA	2022		
198	Glu Lys Ala Asp Ser Cys Glu Lys Glu Ala Leu Met Ser Glu Leu Lys			
199	625 630 635			
200	ATG ATG ACC CAC CTG GGA CAC CAT GAC AAC ATC GTG AAT CTG CTG GGG	2070		
201	Met Met Thr His Leu Gly His His Asp Asn Ile Val Asn Leu Leu Gly			
202	640 645 650			
203	GCA TGC ACA CTG TCA GGG CCA GTG TAC TTG ATT TTT GAA TAT TGT TGC	2118		
204	Ala Cys Thr Leu Ser Gly Pro Val Tyr Leu Ile Phe Glu Tyr Cys Cys			
205	655 660 665			
206	TAT GGT GAC CTC CTC AAC TAC CTA AGA AGT AAA AGA GAG AAG TTT CAC	2166		
207	Tyr Gly Asp Leu Leu Asn Tyr Leu Arg Ser Lys Arg Glu Lys Phe His			
208	670 675 680 685			
209	AGG ACA TGG ACA GAG ATT TTT AAG GAA CAT AAT TTC AGT TCT TAC CCT	2214		
210	Arg Thr Trp Thr Glu Ile Phe Lys Glu His Asn Phe Ser Ser Tyr Pro			
211	690 695 700			
212	ACT TTC CAG GCA CAT TCA AAT TCC AGC ATG CCT GGT TCA CGA GAA GTT	2262		
213	Thr Phe Gln Ala His Ser Asn Ser Ser Met Pro Gly Ser Arg Glu Val			
214	705 710 715			
215	CAG TTA CAC CCG CCC TTG GAT CAG CTC TCA GGG TTC AAT GGG AAT TCA	2310		
216	Gln Leu His Pro Pro Leu Asp Gln Leu Ser Gly Phe Asn Gly Asn Ser			
217	720 725 730			
218	ATT CAT TCT GAA GAT GAG ATT GAA TAT GAA AAC CAG AAG AGG CTG GCA	2358		
219	Ile His Ser Glu Asp Glu Ile Glu Tyr Glu Asn Gln Lys Arg Leu Ala			
220	735 740 745			
221	GAA GAA GAG GAG GAA GAT TTG AAC GTG CTG ACG TTT GAA GAC CTC CTT	2406		
222	Glu Glu Glu Glu Glu Asp Leu Asn Val Leu Thr Phe Glu Asp Leu Leu			
223	750 755 760 765			
224	TGC TTT GCG TAC CAA GTG GCC AAA GGC ATG GAA TTC CTG GAG TTC AAG	2454		
225	Cys Phe Ala Tyr Gln Val Ala Lys Gly Met Glu Phe Leu Glu Phe Lys			
226	770 775 780			
227	TCG TGT GTC CAC AGA GAC CTG GCA GCC AGG AAT GTG TTG GTC ACC CAC	2502		
228	Ser Cys Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Thr His			
229	785 790 795			
230	GGG AAG GTG GTG AAG ATC TGT GAC TTT GGA CTG GCC CGA GAC ATC CTG	2550		
231	Gly Lys Val Val Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ile Leu			
232	800 805 810			
233	AGC GAC TCC AGC TAC GTC GTC AGG GGC AAC GCA CGG CTG CCG GTG AAG	2598		
234	Ser Asp Ser Ser Tyr Val Val Arg Gly Asn Ala Arg Leu Pro Val Lys			
235	815 820 825			
236	TGG ATG GCA CCC GAG AGC TTA TTT GAA GGG ATC TAC ACA ATC AAG AGT	2646		
237	Trp Met Ala Pro Glu Ser Leu Phe Glu Gly Ile Tyr Thr Ile Lys Ser			
238	830 835 840 845			

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : N:\Crf4\05272003\I919408A.raw
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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:5; Line(s) 1031

VERIFICATION SUMMARY

DATE: 05/28/2003

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Input Set : N:\Crf4\05272003\I919408A.raw

Output Set: N:\CRF4\05282003\I919408A.raw

L:20 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:21 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:25 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:27 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:29 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:31 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:33 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:35 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:37 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:39 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:41 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:43 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)
L:45 M:238 W: Alpha Fields not Ordered, Reordered [(A) APPLICATION NUMBER:] of (1) (vii)